

# SEQUENCE LISTING

<110> Dulac, Catherine  
Axel, Richard

<120> Cloning Of Vertebrate Pheromone Receptors And Uses  
Thereof

<130> 0575/48557A/JPW/ADM

<140>

<141>

<150> 08/731,745

<151> 1996-10-18

<160> 21

<170> PatentIn Ver. 2.1

<210> 1

<211> 530

<212> DNA

<213> Homo sapiens

<400> 1

```
aaacataagt ccagttatct acaggtacag gttgatgaga ggccctctcca ttccaccac 60
ctgcctgttg agtgtcctcc aggccatcaa cctcacccca aggagctccc gtttggcaat 120
gttcagagat cctcacatca caaacgcgt tgctttctct tgetgtgggt ctccacata 180
tccattagtg gaagcttctt agtctccact cttccctcca aaaatgttgc ctcaaatagt 240
gttacatttg tcaactcaatc ctgctctgct gggccctga gttgcttct tgggcagaca 300
attttcacac tgatgacatt tcaggatgtc tcttgcagc tcatggcccc cttcagtgga 360
tacatgggga ttctcttggt caggcataac aggcagtctc agcatcttca tagtatcaac 420
ctttctccaa aagcaccccc agataaaagg gccatccaga gcattctttt gctcgtgagt 480
ttctttgtgt tcatgtgcct tttccattt gctgccttaa cacttctgtc 530
```

<210> 2

<211> 1385

<212> DNA

<213> Rattus sp.

<400> 2

```
tttcggcacg agttcacctg ccctcgaatt tcaatttgag taagtgacca gcaatggagt 60
acagaatcag aagatgggtg gatcccaggc aggtgtgtgg aggaggaact ctggaactgc 120
atgaggagtt tgagcacctg ccatggagta gctgatctct gaggaccct cacacaggtc 180
ctgtgtttta catcaagtgc atatttttcc taggatattc atttcgtaa gtctgaaat 240
tacttaattt ttataggagt tctcatatat gatgaataag aacagcagac totacactga 300
```

ttctaacata	aggaataacct	ttttcogctga	aattggcatt	ggagtctcag	ccaatagcct	360
cctactttctc	ttcaacatct	tcaagttaat	ttgtgggcag	aggtccagac	tcactgacct	420
gccattgggt	ctcttgtccc	taatcaactt	acttatgcta	ctgatgacgg	cattcatagc	480
cacagacact	tttattttctt	ggagaggggtg	ggatgacatc	atatgtaaat	cccttctcta	540
cctgtacaga	actttttagag	gtctctctct	ttgtaccagc	tgctgttga	gtgtcctgca	600
ggccatcatc	ctcagtccca	gaagctcctg	tttagcaaag	ttcaaacata	agccttccca	660
tcacatctcc	tgtgccattc	tttctctgag	tgctctctac	atgttcatta	gcagtccacct	720
cttagtatcc	atcattgccca	ccccaaattt	gaccacgaat	gactttattc	atgttactca	780
gtgggtgctct	attctaccca	tgagttacct	catgcaaagc	atgttttcta	cactgctggc	840
catcagggat	gtcttttotta	ttagtctcat	ggctctgtca	acatggtaca	tggtggctct	900
cttgtgtagg	cacaggaaac	agaccoggca	tcttcagggt	accagccttt	ccccaaaagc	960
atcccagaa	caaagggccca	cccgttccat	cctgatgctc	atgagcttat	ttgttctgat	1020
gtctgtcttt	gacagcattg	tctgcagctc	aagaactatg	tatctgaatg	atccaatatc	1080
ttattcttat	caactattta	tgggtgcacat	ctatgccaca	gtaagccctt	ttgtgtttat	1140
tgtcactgaa	aaacatatag	ttaactcttt	gaggtccatg	tgtgtgaagg	tgcataatg	1200
tttgaatatt	cottgatagc	aagctccatt	aagaggagcc	aatgtaagca	tcagaactgt	1260
caatcatggc	gtgctatgtg	ctttggcata	tgtgaaatat	gaagttgttt	ttctgttaaa	1320
atgatttact	ttaactgacg	agatgatgaa	cgtaacagaa	gattaaacca	catccccctt	1380
gatat						1385

<210> 3

<211> 1331

<212> DNA

<213> Rattus sp.

<400> 3

gtggatcccc	cgggctgcag	gaattcggca	cgagccgtga	ttaagggact	ttgaactttt	60
caagggattt	ggagttttat	gaagaatttg	aagatttaca	gagtttacag	gaatggagct	120
gaccagccac	tatgacatgc	cttatatctc	caagagcata	aatataaggc	atggcatgag	180
aggaccagca	gccactgttc	tcatatatga	tgaataagaa	cagcagagtc	cacactgatt	240
ctaccataag	gaataacctc	tccactgaaa	ttggcattgg	aatcttagcc	aacagtttcc	300
tactttctct	ccacatcttc	aagttttattc	gtggacagag	gtccagactc	actgacctgc	360
ccattgggtct	cttgtcccta	atccacctac	tgatgctact	gatgggggca	ttcatagcca	420
tagacatttt	tatttcttgg	aggggatggg	atgacatcat	atgtaaattc	cttgtctact	480
tgtacagaag	tttttagagg	ctctctcttt	gtaccacctg	catgttgagt	gtcctgcagg	540
ccatcaccct	cagccccaga	agctcctggt	tagcaaagtt	caaacataag	tctccccatc	600
acgtctcctg	tgccattatt	tcgctgagca	tcctctacat	gttcattagc	agtcacctct	660
tagtatccat	caatgccacc	cccaatttga	ccacgaacaa	ctttatgcaa	gttactcagt	720
cctgtctacat	tatacccttg	agttacctca	tgcaaagcat	gttttctaca	cttctggcca	780
tcagagatat	ctctcttatt	agtctcatgg	tcctctcgac	ttgttacatg	gaggttctct	840
tgtgtaggca	caggaatcag	atccagcatc	ttcaagggac	caacctttcc	ccaaaagcat	900
ctccagaaca	aagggccaca	cagaccatcc	tgatgctcat	gaccttcttt	gtcctaattgt	960
ccattttcga	cagcattgtc	tcctgttcaa	gaactatgta	tctgaatgat	ccaacatctt	1020
actatattca	aatatttgta	gtggacatct	atgccacagt	cagccctttt	gtgtttatga	1080
gcactggaaa	acatatagtt	aactttttga	agtccatgtg	tgtgaggggtg	aagaatgttt	1140
gaatattcat	taatggacaa	gatcctttaa	gaggagccaa	tgtagtcatc	agaactgtca	1200
gtcatgggtgt	gctgtctatg	tgctttggta	aatgtgaatc	atgaagttgt	ttttctggta	1260

aaatgattta cttaaccaa ctcatgattg taaacatgta acaggagatt aaacaatatc 1320  
cccttcggaa a 1331

<210> 4  
<211> 1496  
<212> DNA  
<213> Rattus sp.

<400> 4  
aattcggcac gagcaaaggc aggggaagatg ctccactggg atgtcatgtc tctatgctcc 60  
acagtggaaa agttgtcaca ttgtacaaac actaaaatta cgaattgctc acaggcacta 120  
aaagcttcct taatcctgtg caggatctcc tcagggtacag agtcctcctg atacgtctat 180  
ctggtcagag gaaagagctg atcagtcatt aacagagctg atttggtccc tccaagggtca 240  
catgacaagg actgtatgag aaaaccagca gtgacatgtc tatagagatc attctgtgcc 300  
acaccagct ccatgtttgg ttgtgtgtat ttgcttccta tccacatata atgaataaag 360  
acaacacact ccatgttgac acaatcatga aaatcactat gttctctgaa gtgagtgttg 420  
gcatcttagc taacagtatc ctgttttttg gtcacctgtg catgtccctt ggagagaaca 480  
agcctaagcc cattcatctc tacattgcat ccttgctcct aacacaacta atgtgtctta 540  
taactatggg actcatagct gctgacatgt ttatttctca ggggatatgg gattctacct 600  
catgccagtc ccttatctat ttgcacaggc ttctgagggg ttttaccctt agtctgcct 660  
gtctgtgaa tgtcttttg atgatactc tcagttctaa aaaatcctgt ttaacaaagt 720  
ttaaacataa ctctcccat cactctcag gtgcctttct tctcctctgt gttctctaca 780  
tgtgttttag cagtcacctt attttatcga ttattgctac ccctaacttg acctcagata 840  
attttatgta tgttactaag tctgttcat ttctacccat gtgttactcc agaacaagca 900  
tgttttccac aacaattgtc gtcagggaag cttttttat cggctctcat gccctgtcca 960  
gtgggtacct ggtggcttct ctctggagac acaggaagca ggcccagcat cttcacagca 1020  
ccggccttct ttc aaagtca tctccagagc aaagggccac cgagaccatc ctgctgtcta 1080  
tgagtttctt tgtgggtctc tacatttttg aaaatgttgt cttctactca aggatgaagt 1140  
tcaaggatgg gtcaacattc tactgtgtcc aaattattgt gtcccatagc tatgccactg 1200  
tcagctcttt tgtgtttatt ttcactgaaa agcgtatgac taagatattg aggtcagtgt 1260  
gtgccagaat aataaataat tgattattca gtgatgggta ttgcccctta gaataaacca 1320  
ttacgtttgc atcagagggt tgggtcatga cataattggg acattctctg tcttaaattg 1380  
ataaatgaaa ttttcttttt tctgtttaa actgtttcct ttgtgtgtgg atgcccata 1440  
tatgaaagaa aactaaacac catgtcctct tacatatcca accaaaaaaa aaaaaa 1496

<210> 5  
<211> 1053  
<212> DNA  
<213> Rattus sp.

<400> 5  
ttttttccca cctcttcatg ctctttgaaa agaacagatc taagcccatt gatctctaca 60  
ttgctttctt atccttaacc caactaatgc tgcttataac tattggactt atagctgcag 120  
acatgtttat gtctcggggg agatgggatt ctaccacatg ccagtccctt atctatttgg 180  
acaggctttt gaggggtttt accctttgtg ctacctgtct gctgaatgtc ctttggacca 240  
tcactctcag tctagaagc tctgttttaa caacatttaa acataaatct ccccatcaca 300

tctcaggtgc	ctttcttttc	ttctgtgttc	tctatatata	ttttggcagt	cacctctttt	360
tatcaacaat	tgctaccccc	aatttgactt	cagataaatt	tatgtatgtt	actaaatcct	420
gttcattttct	acccatgagt	tactccagaa	caagcatgtt	ttccacacca	atggccatca	480
gggaagccct	tcttattggg	ctcattggcc	tgctcagtgg	gtacatgggt	gctttcctat	540
ggagacacaa	gaatcaggcc	cggcatcttc	acagcaccag	cctttcttca	aaagtgtccc	600
cagagcaaaag	ggccaccagg	accatcatga	ttctcatgag	cttctttgtg	gttctctaca	660
ttttggaaaa	tggtgtcttc	tactctagga	tgacattcaa	ggatgggtca	atgttctact	720
gtgtccaaat	tattgtgtcc	catagctatg	ccaccatcag	cccttttgtg	tttatttgca	780
cagaaaagcg	tataattaaa	ctttgggggt	caatgtctag	cagaatagta	agtatttgat	840
tactcagtga	tggatatggg	cccttaatat	aaaccaatat	gttgtcataa	taactatgga	900
tcatgacata	ttggggacat	tctgtgtctt	aaatttataa	aaaaaatttt	ctttttttgt	960
gtttaatctg	tttcccttgt	gtgtggatga	taagtatata	aagggaaatt	aaacagcgtg	1020
tcccctcaga	tatccaaaaa	aaaaaaaaaa	aaa			1053

<210> 6

<211> 1538

<212> DNA

<213> Rattus sp.

<400> 6

gggctgcagg	aattcggcac	gagtcagagt	ccttccttgc	tatgtgtatc	tggagccagc	60
gactcttcta	tggagagcag	ctgtgcaggc	aggtgggtga	gcggaagaag	gcgtgctgct	120
gtgacatcat	caagatgctg	cctagccctg	cgtcgctgct	cttctgagga	agcaggagac	180
tgacccctgt	gacaatgact	tgatgagtc	ctctgttgtc	tacttaccct	agttctttgt	240
cccatacaat	gaggagaatc	agcacactgt	atggagttgt	tgacaagcaa	gctatatttt	300
tctctgaagt	agtcacggg	atctcattca	acagtatcct	cttctcttcc	cacatctttc	360
agttccttct	tgagcgtagg	ctccggatca	ctgacctgat	catcagtctc	ttggccctca	420
tccaccttgg	gatgctaaca	gtcatgggat	tcagagctgt	tgataatttt	gcattctcaga	480
atgtgtggaa	tgacatcaaa	tgcaaatccc	ttgcccaatt	acacagactt	ttgagggggc	540
tctctctttg	tgctacctgt	ctgctgagta	tcttcagggc	catcaccctt	agccccagaa	600
gctcctgttt	agcaaagtcc	aaatataaat	ccacacagca	cagcctgtgt	tcccttcttg	660
tgtcttgggc	cttctacatg	tcctgtggta	ctcactactc	cttcaccatc	gttctgtgact	720
acaacttctc	ttcacgcagt	ctcatatttg	tcactgaatc	ctgcattatt	ttaccatggg	780
attacatcac	cagggattta	tttttcatat	tggggatatt	tcgggatgtg	tccttcatag	840
gtctcatggc	cctctccagc	gggtacatgg	tggccctctt	gtgcagacac	aggaaacagg	900
cccagcatct	tcacaggacc	agcctttctc	caaaagcatc	cccagagcaa	aggggccacca	960
ggaccatcct	gttgctcatg	agcttctttg	tggtgatgta	ctgcttggac	tgaccatat	1020
ccgcctccag	acttatgcac	aacgggtgaac	caatccacca	cagtattcag	atgatgggtct	1080
ccaatagcta	tgccaccctc	agccctttgc	tgtaatttgt	tactgaaaat	ogaattagta	1140
ggtttttgaa	gtccttgcta	ggaaggacag	tagatgctta	agtattgagg	ggaggcaggc	1200
ccactaaagg	agccaatatg	ctagctactg	aataatgaat	cctggcctag	tcctcatgca	1260
atcctgaaca	aattaataca	tgactcatgc	ttcgttaaac	ctgcttcttt	tgaaatgtgt	1320
attaccaaca	cctgtagata	tttgagtcaa	atttcttcat	gtgtattttct	tctcagtgtc	1380
agtaggggac	atctgtgaca	ctttcacaga	ttagggtaac	ttgtgcactt	atcaataagc	1440
taaagtgtac	agcacatttt	actaagccaa	ttatctcaac	agtttgtttt	ctaccaatt	1500
aaatatgtaa	atgttaccac	caaaaaaaaa	aaaaaaaaa			1538

<210> 7  
 <211> 1264  
 <212> DNA  
 <213> Rattus sp.

<400> 7  
 ttggggtaaa acggctcgat gacttccaca tgttttgcca tggcagaatc tgctccatgc 60  
 gggacaagaa aatctctttt ctggtctgac gggcttactg ctgaattcac tgcgcggcga 120  
 ggtaagttga tgactcatga tgaacctgt tctatggctc cagatgacaa acatgatctc 180  
 atatcaggga cttgttcgca ccttccctaa cagtatcctg ttttttgccc acctctgcat 240  
 gttctttgaa gagaacagggt ctaagcccat tgatctgtgc attgctttct tacccttaac 300  
 ccaactaatg ctgcttgtaa ctatgggact catagctgca gacatgttta tggctcaggg 360  
 gatatgggat attaccacat gcaggctcct tatctatctt cacagacttt tgagggggtt 420  
 caacctttgt gctgcctgtc tactgcatac cttttggacc ttcactctca gtcctagaag 480  
 ctctgttta acaaagttta aacataaata tccccatcac atctcagggt cctatctttt 540  
 ctctgtgtt ctctatatgt ccttttagcag tcacctcttt gtattggtca ttgctacctc 600  
 caatttaacc tcagatcatt ttatgtatgt tactcagtc tgctcacttc taacctgag 660  
 ttactccaga acaagcacgt tttccttact gatggtcacc aggggaagtct ttcttatcag 720  
 tctcatggcc ctgtccagtg ggtacatggg gactctccta tggaggcaca agaagcaggc 780  
 ccagcatctt cacagcacca gactttcttc aaaagcatcc ccacagcaaa gggccaccag 840  
 gaccatcctg ctgcttatga ccttctttgt ggttttctac attttaggca ctggttatctt 900  
 ccactcaagg actaagttca aggatgggtc aatcttctac tgtgtccaaa ttattgtgtc 960  
 ccatagctat gccactatca gccattttgt gtttgtttt totgaaaagc gcataatcaa 1020  
 gttttttaga tcaatgtgtg gcagaatagt aaatacttga ttattcactg atgagtatgg 1080  
 gtcataaata tagtctagta aattgtgatc agagtattgg ctcatgacat attaaaaaca 1140  
 ttctcraatt taagtttaac atataaaatt atcttatttc tcttaaagtgt gtttactttg 1200  
 tgtgtattaa aagtatgtaa aagataatta atccccaat acacctttt ttcaaattaa 1260  
 aaaa 1264

<210> 8  
 <211> 315  
 <212> PRT  
 <213> Rattus sp.

<400> 8  
 Met Met Asn Lys Asn Ser Arg Leu Tyr Thr Asp Ser Asn Ile Arg Asn  
 1 5 10 15  
 Thr Phe Phe Ala Glu Ile Gly Ile Gly Val Ser Ala Asn Ser Leu Leu  
 20 25 30  
 Leu Leu Phe Asn Ile Phe Lys Leu Ile Cys Gly Gln Arg Ser Arg Leu  
 35 40 45  
 Thr Asp Leu Pro Ile Gly Leu Leu Ser Leu Ile Asn Leu Leu Met Leu  
 50 55 60

Leu Met Thr Ala Phe Ile Ala Thr Asp Thr Phe Ile Ser Trp Arg Gly  
65 70 75 80

Trp Asp Asp Ile Ile Cys Lys Ser Leu Leu Tyr Leu Tyr Arg Thr Phe  
85 90 95

Arg Gly Leu Ser Leu Cys Thr Ser Cys Leu Leu Ser Val Leu Gln Ala  
100 105 110

Ile Ile Leu Ser Pro Arg Ser Ser Cys Leu Ala Lys Phe Lys His Lys  
115 120 125

Pro Ser His His Ile Ser Cys Ala Ile Leu Ser Leu Ser Val Leu Tyr  
130 135 140

Met Phe Ile Ser Ser His Leu Leu Val Ser Ile Ile Ala Thr Pro Asn  
145 150 155 160

Leu Thr Thr Asn Asp Phe Ile His Val Thr Gln Trp Cys Ser Ile Leu  
165 170 175

Pro Met Ser Tyr Leu Met Gln Ser Met Phe Ser Thr Leu Leu Ala Ile  
180 185 190

Arg Asp Val Phe Leu Ile Ser Leu Met Val Leu Ser Thr Trp Tyr Met  
195 200 205

Val Ala Leu Leu Cys Arg His Arg Lys Gln Thr Arg His Leu Gln Gly  
210 215 220

Thr Ser Leu Ser Pro Lys Ala Ser Pro Glu Gln Arg Ala Thr Arg Ser  
225 230 235 240

Ile Leu Met Leu Met Ser Leu Phe Val Leu Met Ser Val Phe Asp Ser  
245 250 255

Ile Val Cys Ser Ser Arg Thr Met Tyr Leu Asn Asp Pro Ile Ser Tyr  
260 265 270

Ser Tyr Gln Leu Phe Met Val His Ile Tyr Ala Thr Val Ser Pro Phe  
275 280 285

Val Phe Ile Val Thr Glu Lys His Ile Val Asn Ser Leu Arg Ser Met  
290 295 300

Cys Val Lys Val His Glu Cys Leu Asn Ile Pro  
305 310 315

<210> 9  
 <211> 311  
 <212> PRT  
 <213> Rattus sp.

<400> 9

Met	Met	Asn	Lys	Asn	Ser	Arg	Leu	His	Ile	Asp	Ser	Asn	Ile	Arg	Asn	1	5	10	15
Thr	Phe	Phe	Thr	Glu	Ile	Gly	Ile	Gly	Val	Ser	Ala	Asn	Ser	Leu	Leu	20	25	30	
Leu	Leu	Phe	Asn	Ile	Phe	Lys	Phe	Ile	His	Gly	Gln	Arg	Ser	Arg	Leu	35	40	45	
Thr	Asp	Leu	Pro	Ile	Gly	Leu	Leu	Ser	Leu	Ile	Asn	Leu	Leu	Met	Leu	50	55	60	
Leu	Ile	Met	Ala	Cys	Ile	Ala	Thr	Asp	Ile	Phe	Ile	Ser	Cys	Arg	Arg	65	70	75	80
Trp	Asp	Asp	Ile	Ile	Cys	Lys	Ser	Leu	Leu	Tyr	Leu	Tyr	Arg	Thr	Phe	85	90	95	
Arg	Gly	Leu	Ser	Leu	Ser	Thr	Thr	Cys	Leu	Leu	Ser	Val	Leu	Gln	Ala	100	105	110	
Ile	Ile	Leu	Ser	Pro	Arg	Ser	Ser	Cys	Leu	Ala	Lys	Tyr	Lys	His	Lys	115	120	125	
Pro	Pro	His	His	Ile	Phe	Cys	Ala	Met	Leu	Phe	Leu	Ser	Val	Leu	Tyr	130	135	140	
Met	Phe	Ile	Ser	Ser	His	Leu	Leu	Leu	Ser	Ile	Ile	Ala	Thr	Pro	Asn	145	150	155	160
Leu	Thr	Thr	Asn	Asp	Phe	Ile	His	Val	Ser	Gln	Ser	Cys	Ser	Ile	Leu	165	170	175	
Pro	Met	Ser	Tyr	Leu	Met	Gln	Ser	Met	Phe	Ser	Thr	Leu	Leu	Ala	Ile	180	185	190	
Arg	Asn	Val	Phe	Leu	Ile	Ser	Leu	Ile	Val	Leu	Ser	Thr	Trp	Tyr	Met	195	200	205	
Val	Ala	Leu	Leu	Cys	Arg	His	Arg	Lys	Gln	Thr	Arg	His	Leu	Gln	Asp				





Ile Thr Leu Ser Pro Arg Ser Ser Cys Leu Ala Lys Phe Lys His Lys  
 115 120 125

Ser Pro His His Val Ser Cys Ala Ile Ile Ser Leu Ser Ile Leu Tyr  
 130 135 140

Met Phe Ile Ser Ser His Leu Leu Val Ser Ile Asn Ala Thr Pro Asn  
 145 150 155 160

Leu Thr Thr Asn Asn Phe Met Gln Val Thr Gln Ser Cys Tyr Ile Ile  
 165 170 175

Pro Leu Ser Tyr Leu Met Gln Ser Met Phe Ser Thr Leu Leu Ala Ile  
 180 185 190

Arg Asp Ile Ser Leu Ile Ser Leu Met Val Leu Ser Thr Cys Tyr Met  
 195 200 205

Glu Val Leu Leu Cys Arg His Arg Asn Gln Ile Gln His Leu Gln Gly  
 210 215 220

Thr Asn Leu Ser Pro Lys Ala Ser Pro Glu Gln Arg Ala Thr Gln Thr  
 225 230 235 240

Ile Leu Met Leu Met Thr Phe Phe Val Leu Met Ser Ile Phe Asp Ser  
 245 250 255

Ile Val Ser Cys Ser Arg Thr Met Tyr Leu Asn Asp Pro Thr Ser Tyr  
 260 265 270

Tyr Ile Gln Ile Phe Gly Val Asp Ile Tyr Ala Thr Val Ser Pro Phe  
 275 280 285

Val Phe Met Ser Thr Glu Lys His Ile Val Asn Phe Leu Lys Ser Met  
 290 295 300

Cys Val Arg Val Lys Asn Val  
 305 310

<210> 11  
 <211> 310  
 <212> PRT  
 <213> Rattus sp.

<400> 11  
 Met Asn Lys Asp Asn Thr Leu His Val Asp Thr Ile Met Lys Ile Thr  
 1 5 10 15

Met Phe Ser Glu Val Ser Val Gly Ile Leu Ala Asn Ser Ile Leu Phe  
 20 25 30  
 Phe Gly His Leu Cys Met Leu Leu Gly Glu Asn Lys Pro Lys Pro Ile  
 35 40 45  
 His Leu Tyr Ile Ala Ser Leu Ser Leu Thr Gln Leu Met Leu Leu Ile  
 50 55 60  
 Thr Met Gly Leu Ile Ala Ala Asp Met Phe Ile Ser Gln Gly Ile Trp  
 65 70 75 80  
 Asp Ser Thr Ser Cys Gln Ser Leu Ile Tyr Leu His Arg Leu Ser Arg  
 85 90 95  
 Gly Phe Thr Leu Ser Ala Ala Cys Leu Leu Asn Val Phe Trp Met Ile  
 100 105 110  
 Thr Leu Ser Ser Lys Lys Ser Cys Leu Thr Lys Phe Lys His Asn Ser  
 115 120 125  
 Pro His His Ile Ser Gly Ala Phe Leu Leu Leu Cys Val Leu Tyr Met  
 130 135 140  
 Cys Phe Ser Ser His Leu Ile Leu Ser Ile Ile Ala Thr Pro Asn Leu  
 145 150 155 160  
 Thr Ser Asp Asn Phe Met Tyr Val Thr Lys Ser Cys Ser Phe Leu Pro  
 165 170 175  
 Met Cys Tyr Ser Arg Thr Ser Met Phe Ser Thr Thr Ile Ala Val Arg  
 180 185 190  
 Glu Ala Phe Phe Ile Gly Leu Met Ala Leu Ser Ser Gly Tyr Leu Val  
 195 200 205  
 Ala Phe Leu Trp Arg His Arg Lys Gln Ala Gln His Leu His Ser Thr  
 210 215 220  
 Gly Leu Ser Ser Lys Ser Ser Pro Glu Gln Arg Ala Thr Glu Thr Ile  
 225 230 235 240  
 Leu Leu Leu Met Ser Phe Phe Val Val Leu Tyr Ile Leu Glu Asn Val  
 245 250 255  
 Val Phe Tyr Ser Ser Arg Met Phe Lys Asp Gly Ser Thr Phe Tyr Cys  
 260 265 270

Val Gln Ile Ile Val Ser His Ser Tyr Ala Thr Val Ser Ser Phe Val  
 275 280 285

Phe Ile Phe Thr Glu Lys Arg Met Thr Lys Ile Leu Arg Ser Val Cys  
 290 295 300

Ala Arg Ile Ile Asn Asn  
 305 310

<210> 12  
 <211> 278  
 <212> PRT  
 <213> Rattus sp.

<400> 12  
 Phe Ser His Leu Phe Met Leu Phe Glu Lys Asn Arg Ser Lys Pro Ile  
 1 5 10 15

Asp Leu Tyr Ile Ala Phe Leu Ser Leu Thr Gln Leu Met Leu Leu Ile  
 20 25 30

Thr Ile Gly Leu Ile Ala Ala Asp Met Phe Met Ser Arg Gly Arg Trp  
 35 40 45

Asp Ser Thr Thr Cys Gln Ser Leu Ile Tyr Leu Asp Arg Leu Leu Arg  
 50 55 60

Gly Phe Thr Leu Cys Ala Thr Cys Leu Leu Asn Val Leu Trp Thr Ile  
 65 70 75 80

Thr Leu Ser Pro Arg Ser Ser Cys Leu Thr Thr Phe Lys His Lys Ser  
 85 90 95

Pro His His Ile Ser Gly Ala Phe Leu Phe Phe Cys Val Leu Tyr Ile  
 100 105 110

Ser Phe Gly Ser His Leu Phe Leu Ser Thr Ile Ala Thr Pro Asn Leu  
 115 120 125

Thr Ser Asp Asn Phe Met Tyr Val Thr Lys Ser Cys Ser Phe Leu Pro  
 130 135 140

Met Ser Tyr Ser Arg Thr Ser Met Phe Ser Thr Pro Met Ala Ile Arg  
 145 150 155 160

Glu Ala Leu Leu Ile Gly Leu Ile Gly Leu Ser Ser Gly Tyr Met Val



Gly Leu Ser Leu Cys Ala Thr Cys Leu Leu Ser Ile Phe Gln Ala Ile  
 100 105 110

Thr Leu Ser Pro Arg Ser Ser Cys Leu Ala Lys Phe Lys Tyr Lys Ser  
 115 120 125

Thr Gln His Ser Leu Cys Ser Leu Leu Val Leu Trp Ala Phe Tyr Met  
 130 135 140

Ser Cys Gly Thr His Tyr Ser Phe Thr Ile Val Ala Asp Tyr Asn Phe  
 145 150 155 160

Ser Ser Arg Ser Leu Ile Phe Val Thr Glu Ser Cys Ile Ile Leu Pro  
 165 170 175

Met Asp Tyr Ile Thr Arg His Leu Phe Phe Ile Leu Gly Ile Phe Arg  
 180 185 190

Asp Val Ser Phe Ile Gly Leu Met Ala Leu Ser Ser Gly Tyr Met Val  
 195 200 205

Ala Leu Leu Cys Arg His Arg Lys Gln Ala Gln His Leu His Arg Thr  
 210 215 220

Ser Leu Ser Pro Lys Ala Ser Pro Glu Gln Arg Ala Thr Arg Thr Ile  
 225 230 235 240

Leu Leu Leu Met Ser Phe Phe Val Leu Met Tyr Cys Leu Asp Cys Thr  
 245 250 255

Ile Ser Ala Ser Arg Leu Met His Asn Gly Glu Pro Ile His His Ser  
 260 265 270

Ile Gln Met Met Val Ser Asn Ser Tyr Ala Thr Leu Ser Pro Leu Leu  
 275 280 285

Leu Ile Val Thr Glu Asn Arg Ile Ser Arg Phe Leu Lys Ser Leu Leu  
 290 295 300

Gly Arg Thr Val Asp Ala  
 305 310

<210> 14  
 <211> 307  
 <212> PRT  
 <213> Rattus sp.

<400> 14

Met Met Asn Pro Val Leu Trp Leu Gln Met Thr Asn Met Ile Ser Tyr  
1 5 10 15

Gln Gly Leu Val Arg Thr Phe Pro Asn Ser Ile Leu Phe Phe Ala His  
20 25 30

Leu Cys Met Phe Phe Glu Glu Asn Arg Ser Lys Pro Ile Asp Leu Cys  
35 40 45

Ile Ala Phe Leu Ser Leu Thr Gln Leu Met Leu Leu Val Thr Met Gly  
50 55 60

Leu Ile Ala Ala Asp Met Phe Met Ala Gln Gly Ile Trp Asp Ile Thr  
65 70 75 80

Thr Cys Arg Ser Leu Ile Tyr Phe His Arg Leu Leu Arg Gly Phe Asn  
85 90 95

Leu Cys Ala Ala Cys Leu Leu His Ile Leu Trp Thr Phe Thr Leu Ser  
100 105 110

Pro Arg Ser Ser Cys Leu Thr Lys Phe Lys His Lys Ser Pro His His  
115 120 125

Ile Ser Gly Ala Tyr Leu Phe Phe Cys Val Leu Tyr Met Ser Phe Ser  
130 135 140

Ser His Leu Phe Val Leu Val Ile Ala Thr Ser Asn Leu Thr Ser Asp  
145 150 155 160

His Phe Met Tyr Val Thr Gln Ser Cys Ser Leu Leu Pro Met Ser Tyr  
165 170 175

Ser Arg Thr Ser Thr Phe Ser Leu Leu Met Val Thr Arg Glu Val Phe  
180 185 190

Leu Ile Ser Leu Met Ala Leu Ser Ser Gly Tyr Met Val Thr Leu Leu  
195 200 205

Trp Arg His Lys Lys Gln Ala Gln His Leu His Ser Thr Arg Leu Ser  
210 215 220

Ser Lys Ala Ser Pro Gln Gln Arg Ala Thr Arg Thr Ile Leu Leu Leu  
225 230 235 240

Met Thr Phe Phe Val Val Phe Tyr Ile Leu Gly Thr Val Ile Phe His  
245 250 255

Ser Arg Thr Lys Phe Lys Asp Gly Ser Ile Phe Tyr Cys Val Gln Ile  
260 265 270

Ile Val Ser His Ser Tyr Ala Thr Ile Ser Pro Phe Val Phe Val Phe  
275 280 285

Ser Glu Lys Arg Ile Ile Lys Phe Phe Arg Ser Met Cys Gly Arg Ile  
290 295 300

Val Asn Thr  
305

<210> 15  
<211> 173  
<212> PRT  
<213> Homo sapiens

<400> 15  
Asn Ile Ser Pro Val Ile Tyr Arg Tyr Arg Leu Met Arg Gly Leu Ser  
1 5 10 15

Ile Ser Thr Thr Cys Leu Leu Ser Val Leu Gln Ala Ile Asn Leu Thr  
20 25 30

Pro Arg Ser Ser Arg Leu Ala Arg Ser Ser His His Lys Pro Arg Cys  
35 40 45

Phe Leu Leu Leu Trp Val Phe His Ile Ser Ile Ser Gly Ser Phe Leu  
50 55 60

Val Ser Thr Leu Pro Ser Lys Asn Val Ala Ser Asn Ser Val Thr Phe  
65 70 75 80

Val Thr Gln Ser Cys Ser Ala Gly Pro Leu Ser Cys Phe Leu Gly Gln  
85 90 95

Thr Ile Phe Thr Leu Met Thr Phe Gln Asp Val Ser Leu Gln Leu Met  
100 105 110

Ala Pro Phe Ser Gly Tyr Met Val Ile Leu Leu Cys Arg His Asn Arg  
115 120 125

Gln Ser Gln His Leu His Ser Ile Asn Leu Ser Pro Lys Ala Pro Pro  
130 135 140

Asp Lys Arg Ala Ile Gln Ser Ile Leu Leu Leu Val Ser Phe Phe Val

145 150 155 160

Phe Met Cys Leu Phe Pro Phe Ala Ala Leu Thr Leu Leu  
165 170

<210> 16  
<211> 71  
<212> PRT  
<213> Rattus sp.

<400> 16  
Ser Lys Arg Lys Lys Ser Phe Leu Leu Cys Ile Gly Trp Leu Ala Leu  
1 5 10 15

Thr Asp Leu Val Gly Gln Leu Leu Thr Ser Pro Val Val Ile Leu Val  
20 25 30

Tyr Leu Ser Gln Arg Arg Trp Glu Gln Leu Asp Pro Ser Gly Arg Leu  
35 40 45

Cys Thr Phe Phe Gly Leu Thr Met Thr Val Phe Gly Leu Ser Ser Leu  
50 55 60

Leu Val Ala Ser Ala Met Ala  
65 70

<210> 17  
<211> 74  
<212> PRT  
<213> Rattus sp.

<400> 17  
Gly Gln Arg Ser Arg Leu Thr Asp Leu Pro Ile Gly Leu Leu Ser Leu  
1 5 10 15

Ile Asn Leu Leu Met Leu Leu Ile Met Ala Cys Ile Ala Thr Asp Ile  
20 25 30

Phe Ile Ser Cys Arg Arg Trp Asp Asp Ile Ile Cys Lys Ser Leu Leu  
35 40 45

Tyr Leu Tyr Arg Thr Phe Arg Gly Leu Ser Leu Ser Thr Thr Cys Leu  
50 55 60

Leu Ser Val Leu Gln Ala Ile Ile Leu Ser  
65 70



<210> 18  
 <211> 174  
 <212> PRT  
 <213> Rattus sp.

<400> 18  
 Lys Cys Lys Ser Leu Ala His Leu His Arg Leu Leu Arg Gly Leu Ser  
 1 5 10 15

Leu Cys Ala Thr Cys Leu Leu Ser Ile Phe Gln Ala Ile Thr Leu Ser  
 20 25 30

Pro Arg Ser Ser Cys Leu Ala Lys Ser Thr Gln His Ser Leu Cys Ser  
 35 40 45

Leu Leu Val Leu Trp Ala Phe Tyr Met Ser Cys Gly Thr His Tyr Ser  
 50 55 60

Phe Thr Ile Val Ala Asp Tyr Asn Phe Ser Ser Arg Ser Leu Ile Phe  
 65 70 75 80

Val Thr Glu Ser Cys Ile Ile Leu Pro Met Asp Tyr Ile Thr Arg Asp  
 85 90 95

Leu Phe Phe Ile Leu Gly Ile Phe Arg Asp Val Ser Phe Ile Gly Leu  
 100 105 110

Met Ala Leu Ser Ser Gly Tyr Met Val Ala Leu Leu Cys Arg His Arg  
 115 120 125

Lys Gly Ala Gln His Leu His Arg Thr Ser Leu Ser Pro Lys Ala Ser  
 130 135 140

Pro Glu Gln Arg Ala Thr Arg Thr Ile Leu Leu Leu Met Ser Phe Phe  
 145 150 155 160

Val Leu Met Tyr Cys Leu Asp Cys Thr Ile Ser Ala Ser Arg  
 165 170

<210> 19  
 <211> 32  
 <212> PRT  
 <213> Rattus sp.

<220>

<221> VARIANT  
 <222> (3)  
 <223> Xaa at position 3 is Leu or Phe  
  
 <220>  
 <221> VARIANT  
 <222> (4)  
 <223> Xaa at position 4 is Ser or Thr or Asn  
  
 <220>  
 <221> VARIANT  
 <222> (6)  
 <223> Xaa at position 6 is Cys or Ser  
  
 <220>  
 <221> VARIANT  
 <222> (7)  
 <223> Xaa at position 7 is Ala or Thr  
  
 <220>  
 <221> VARIANT  
 <222> (8)  
 <223> Xaa at position 8 is Thr or Ala or Ser  
  
 <220>  
 <221> VARIANT  
 <222> (10)  
 <223> Xaa at position 10 is Leu or Met  
  
 <220>  
 <221> VARIANT  
 <222> (12)  
 <223> Xaa at position 12 is Ser or Asn or His  
  
 <220>  
 <221> VARIANT  
 <222> (13)  
 <223> Xaa at position 13 is Val or Ile  
  
 <220>  
 <221> VARIANT  
 <222> (14)  
 <223> Xaa at position 14 is Leu or Phe  
  
 <220>  
 <221> VARIANT  
 <222> (15)  
 <223> Xaa at position 15 is Gln or Trp

<220>  
<221> VARIANT  
<222> (16)  
<223> Xaa at position 16 is Ala or Thr or Met

<220>  
<221> VARIANT  
<222> (17)  
<223> Xaa at position 17 is Ile or Phe

<220>  
<221> VARIANT  
<222> (18)  
<223> Xaa at position 18 is Ile or Thr

<220>  
<221> VARIANT  
<222> (21)  
<223> Xaa at position 21 is Pro or Ser

<220>  
<221> VARIANT  
<222> (22)  
<223> Xaa at position 22 is Arg or Lys

<220>  
<221> VARIANT  
<222> (23)  
<223> Xaa at position 23 is Ser or Lys

<220>  
<221> VARIANT  
<222> (27)  
<223> Xaa at position 27 is Ala or Thr

<220>  
<221> VARIANT  
<222> (28)  
<223> Xaa at position 28 is Lys or Thr

<220>  
<221> VARIANT  
<222> (29)  
<223> Xaa at position 29 is Phe or Tyr

<220>  
<221> VARIANT

<222> (31)  
 <223> Xaa at position 31 is His or Tyr  
  
 <220>  
 <221> VARIANT  
 <222> (32)  
 <223> Xaa at position 32 is Lys or Asn  
  
 <400> 19  
 Arg Gly Xaa Xaa Leu Xaa Xaa Xaa Cys Xaa Leu Xaa Xaa Xaa Xaa Xaa  
 1 5 10 15  
  
 Xaa Xaa Leu Ser Xaa Xaa Xaa Ser Cys Leu Xaa Xaa Xaa Lys Xaa Xaa  
 20 25 30

<210> 20  
 <211> 21  
 <212> PRT  
 <213> Rattus sp.  
  
 <220>  
 <221> VARIANT  
 <222> (2)  
 <223> Xaa at position 2 is Ala or Ser or Val  
  
 <220>  
 <221> VARIANT  
 <222> (5)  
 <223> Xaa at position 5 is Glu or Gln  
  
 <220>  
 <221> VARIANT  
 <222> (10)  
 <223> Xaa at position 10 is Arg or Gln or Glu  
  
 <220>  
 <221> VARIANT  
 <222> (11)  
 <223> Xaa at position 11 is Thr or Ser  
  
 <220>  
 <221> VARIANT  
 <222> (13)  
 <223> Xaa at position 13 is Leu or Met

<220>  
 <221> VARIANT  
 <222> (14)  
 <223> Xaa at position 14 is Met or Leu or Ile

<220>  
 <221> VARIANT  
 <222> (16)  
 <223> Xaa at position 16 is Met or Arg

<220>  
 <221> VARIANT  
 <222> (17)  
 <223> Xaa at position 17 is Ser or Thr

<220>  
 <221> VARIANT  
 <222> (18)  
 <223> Xaa at position 18 is Phe or Leu

<220>  
 <221> VARIANT  
 <222> (20)  
 <223> Xaa at position 20 is Val or Gly

<220>  
 <221> VARIANT  
 <222> (21)  
 <223> Xaa at position 21 is Val or Leu

<400> 20  
 Lys Xaa Ser Pro Xaa Gln Arg Ala Thr Xaa Xaa Ile Xaa Xaa Leu Xaa  
 1 5 10 15

Xaa Xaa Phe Xaa Xaa  
 20

<210> 21  
 <211> 9  
 <212> PRT  
 <213> Rattus sp.

<220>  
 <223> Xaa at position 4 is Val or Ile or Leu, Xaa at  
 position 6 is Pro or Ser, Xaa at position 7 is Phe  
 or Leu, Xaa at position 8 is Val or Leu, Xaa at

position 9 is Phe or Leu

<220>

<221> VARIANT

<222> (4)

<223> Xaa at position 4 is Val or Ile or Leu

<220>

<221> VARIANT

<222> (6)

<223> Xaa at position 6 is Pro or Ser

<220>

<221> VARIANT

<222> (7)

<223> Xaa at position 7 is Phe or Leu

<220>

<221> VARIANT

<222> (8)

<223> Xaa at position 8 is Val or Leu

<220>

<221> VARIANT

<222> (9)

<223> Xaa at position 9 is Phe or Leu

<400> 21

Tyr Ala Thr Xaa Ser Xaa Xaa Xaa Xaa

1

5